

GenCore version 5.1.3
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OM protein - protein search, using sw model

Run on: January 16, 2003, 16:34:37 : Search time 23.7957 seconds
(without alignments)
28.011 Million cell updates/sec

Title: US-09-856-070-25

Perfect score: 23

Sequence: 1 MLPLQ 5

Scoring table: RUSUM62

Gapop 10 0, Gapext 0 5

Searched: 908470 seqs, 13250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0

Maximum DB seq length: 20000000

Post-processing: Minimum Match 68

Maximum Match 100%

Listing first 45 summaries

Database A: Genes-110102 *

1:	/SUS2/accdata/geneseq/geneseq-emb1/AA1980.DAT:
2:	/SUS2/accdata/geneseq/geneseq-emb1/AA1981.DAT:
3:	/SUS2/accdata/geneseq/geneseq-emb1/AA1982.DAT:
4:	/SUS2/accdata/geneseq/geneseq-emb1/AA1983.DAT:
5:	/SUS2/accdata/geneseq/geneseq-emb1/AA1984.DAT:
6:	/SUS2/accdata/geneseq/geneseq-emb1/AA1985.DAT:
7:	/SUS2/accdata/geneseq/geneseq-emb1/AA1986.DAT:
8:	/SUS2/accdata/geneseq/geneseq-emb1/AA1987.DAT:
9:	/SUS2/accdata/geneseq/geneseq-emb1/AA1988.DAT:
10:	/SUS2/accdata/geneseq/geneseq-emb1/AA1989.DAT:
11:	/SUS2/accdata/geneseq/geneseq-emb1/AA1990.DAT:
12:	/SUS2/accdata/geneseq/geneseq-emb1/AA1991.DAT:
13:	/SUS2/accdata/geneseq/geneseq-emb1/AA1992.DAT:
14:	/SUS2/accdata/geneseq/geneseq-emb1/AA1993.DAT:
15:	/SUS2/accdata/geneseq/geneseq-emb1/AA1994.DAT:
16:	/SUS2/accdata/geneseq/geneseq-emb1/AA1995.DAT:
17:	/SUS2/accdata/geneseq/geneseq-emb1/AA1996.DAT:
18:	/SUS2/accdata/geneseq/geneseq-emb1/AA1997.DAT:
19:	/SUS2/accdata/geneseq/geneseq-emb1/AA1998.DAT:
20:	/SUS2/accdata/geneseq/geneseq-emb1/AA1999.DAT:
21:	/SUS2/accdata/geneseq/geneseq-emb1/AA2000.DAT:
22:	/SUS2/accdata/geneseq/geneseq-emb1/AA2001.DAT:
23:	/SUS2/accdata/geneseq/geneseq-emb1/AA2002.DAT:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Match	Score	Length	ID	Description
1	23	100.0	5	AA082040	Human heprecceptor
2	23	100.0	11	AA082039	Human heprecceptor
3	23	100.0	12	AA017959	Integrin binding p
4	23	100.0	12	AA082038	Human heprecceptor
5	23	100.0	12	AA081119	Integrin antagonist
6	23	100.0	12	AA072970	Integrin binding p
7	23	100.0	13	AA082037	Human heprecceptor
8	23	100.0	15	AA082035	Human secreted pro
9	23	100.0	34	AA082020	Human heprecceptor
10	23	100.0	57	AA087003	Human immune/haema

11	23	100.0	73	22	AA056711	Human polypeptide
12	23	100.0	82	22	AB010135	Human cDNA SP, ID
13	23	100.0	109	21	AA053845	Human secreted pro
14	23	100.0	111	22	AA028299	Novel human diapo
15	23	100.0	114	21	AA023000	Arabidopsis thalia
16	23	100.0	114	21	AA047567	Arabidopsis thalia
17	23	100.0	115	21	AA022999	Arabidopsis thalia
18	23	100.0	115	21	AA047566	Arabidopsis thalia
19	23	100.0	116	22	AA074592	Human colon cancer
20	23	100.0	124	21	AA047540	Arabidopsis thalia
21	23	100.0	125	21	AA047539	Arabidopsis thalia
22	23	100.0	134	22	AA067629	Propionibacterium
23	23	100.0	150	23	AB041710	Human ovarian anti
24	23	100.0	166	22	AA060915	Novel human diapo
25	23	100.0	173	21	AA022998	Arabidopsis thalia
26	23	100.0	173	21	AA047565	Arabidopsis thalia
27	23	100.0	183	21	AA047538	Arabidopsis thalia
28	23	100.0	185	21	AA057352	Arabidopsis thalia
29	23	100.0	188	22	AA066050	Propionibacterium
30	23	100.0	166	21	AA057351	Arabidopsis thalia
31	23	100.0	199	23	AB042172	Human ovarian anti
32	23	100.0	204	24	AB019053	Novel human diapo
33	23	100.0	209	21	AA053844	Human secreted pro
34	23	100.0	220	23	AA047545	Listeria monocytog
35	23	100.0	222	21	AA057350	Arabidopsis thalia
36	23	100.0	224	22	AB008709	Novel human diapo
37	23	100.0	239	22	AB011801	Human secreted pro
38	23	100.0	241	22	AA047564	Human protein sequ
39	23	100.0	245	22	AA059320	Novel human diapo
40	24	100.0	272	24	AA048914	Listeria monocytog
41	24	100.0	278	22	AA059803	S-adenosylmethioni
42	24	100.0	309	24	AA073403	Human secreted pro
43	24	100.0	337	24	AA051910	Heptidically activ
44	24	100.0	347	22	AA028300	Novel human diapo
45	24	100.0	448	22	AA069117	Protephila melanog

ALIGNMENTS

RESULT 1
AA082040
1: AA082040 standard, peptide, 5 AA
XX AA082040:
AC AA082040:
13-JUN-2001 (first entry)
Human heprecceptor domain A binding peptide Eupr428.

Human, heprecceptor, cytostatic, anti HIV, antibiotic;
Neutropenic, immune response inducer, cefin, infectious diseases, cancer;
HIV related dementia.

XX Homo sapiens.

XX CB0354241-A.

XX 21-MAR-2001.

XX 17-SEP-1999; 99GB-0021881.

XX 17-SEP-1999; 99GB-0021881.

XX (HOLM/) HOLMS R D.

XX Holms RD;

XX WPI, 2061 293287/31.

XX Novel regulatory or unfolding peptides of cefin that binds to
PI heprecceptor, useful for inducing immune response for treating
PI infectious diseases and cancer.

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XX PS Claim 26: Page 37: 4app: English
XX CC the heprecceptor is a novel active site in human ezrin. Ezrin regulates
XX CC the structure of the cortical cytoskeleton to control cell surface
XX CC topography. The present invention relates to peptides (see AAB82021 to
XX CC AAB82041) that bind to heprecceptor with greater affinity than HEPI (see
XX CC AAB82046). The heprecceptor binding peptides are useful for inducing
XX CC immune response, and for treating infectious diseases, cancer and
XX CC HIV-related dementia. The present peptide binds to domain A of the
XX CC heprecceptor (AAB82019).
XX SQ Sequence 5 AA:

Query Match: 100.0%; Score 23; DB 22; Length 5;
Best local Similarity 100.0%; Pred. No. 19;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MLRLQ 5
DB 1 MLRLQ 5
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DB 3 MLRLQ 7
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RESULT 2
AAB82019
ID AAB82019 standard; peptide; 11 AA.
XX AC AAB82019;
XX AC AAB17969;
XX AC AAB17969;
XX DT 31-OCT-2000 (first entry)
XX DE Integrin-binding peptide sequence SEQ ID No:1081.
XX DE
XX DE
XX KW Modified peptide; therapeutic agent; fusion: FC domain; cancer;
XX KW autoimmune disease; cytostatic; antiasthmatic; thrombolytic; VEGF;
XX KW immunosuppressive; EGF; TPO; CMHA; mimetic; IL-1; TNF; antagonist;
XX KW MMP; inhibitor; erythropoietin; thrombopoietin; interleukin 1;
XX KW cytotoxic T cell lymphocyte antigen 4; tumour necrosis factor;
XX KW vascular endothelial growth factor; matrix metalloproteinase;
XX KW asthma; thrombosis; pharmaceutical.
XX OS Synthetic.
XX OS
XX PN WO200024782-A2.
XX PN
XX PN 04-MAY-2000.
XX PN
XX PN 25-OCT-1999; 99WO-US25044.
XX PR
XX PR 23-OCT-1998; 98US-0105371.
XX PR 22-OCT-1999; 99US-0428082.
XX PR
XX PA (AMGEN) AMGEN INC.
XX PI
XX PI Feige U, Liu C, Cheetham J, Boone TC;
XX PI WPI: 2000-350702/30.
XX PI
XX PI Novel composition of matter comprising an FC domain and
XX PI pharmacologically active peptides, useful for treating cancer and
XX PI autoimmune diseases.
XX PS Claim 39: Page 592: 608pp; English.
XX PS
XX CC The present invention describes composition of matter (1) comprising an
XX CC FC domain, pharmacologically active peptides, and linkers, where (1) is:
XX CC (X1)a-PI-(X2)b, where: P1 = an FC domain; X1 and X2 = are each
XX CC independently selected from -(L1)c-P1, -(L1)c-P1-(L2)d-P2,
XX CC -(L1)c-P1-(L2)d-P2-(L3)e-P3, or -(L1)c-P1-(L2)d-P2-(L3)e-P3-(L4)f-P4
XX CC where P1, P2, P3, and P4 = are each independently sequences of
XX CC pharmacologically active peptides; L1, L2, L3, and L4 = are each
XX CC independently linkers; and a, b, c, d, e, and f = are each independently
XX CC 0 or 1, provided that at least 1 of a and b is 1. The composition can
XX CC have cytostatic, antiasthmatic, thrombolytic and immunosuppressive
XX CC activities. DNAs, vectors and host cells from the present invention can
XX CC be used for producing pharmaceutical compositions. The compositions are
XX CC useful for treating cancer, asthma, thrombosis, or autoimmune diseases.
XX CC The use of an FC domain (rather than a Fab domain) can provide a longer
XX CC half life or incorporate functions such as FC receptor binding, protein
XX CC A binding, complement fixation, and possibly placental transfer. AAB6443
XX CC to AAB6926 and AAB1055 to AAB1863 represent nucleotide and amino acid
XX CC sequences used in the exemplification of the present invention.
XX SQ Sequence 12 AA:

Query Match: 100.0%; Score 23; DB 21; Length 12;
Best local Similarity 100.0%; Pred. No. 19;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MLRLQ 5
DB 1 MLRLQ 5
   | | | | |
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DB 3 MLRLQ 7
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RESULT 2
AAB82019
ID AAB82019 standard; peptide; 11 AA.
XX AC AAB82019;
XX AC
XX DT 13-JUN-2001 (first entry)
XX DE Human heprecceptor domain A binding peptide Rupe2232.
XX DE
XX KW Human; heprecceptor; cytostatic; anti-HIV; antibiotic;
XX KW neutropic; immune response inducer; ezrin, infectious diseases, cancer;
XX KW HIV-related dementia.
XX OS Homo sapiens.
XX OS
XX PN Key Location/Qualifiers
XX PN Modified site 9 /note= "optionally phosphorylated"
XX PN
XX PN GH2354241-A.
XX PN
XX PN 21-MAR-2001.
XX PN
XX PN 17-SEP-1999; 99GB-0021881.
XX PN
XX PN 17-SEP-1999; 99GB-0021881.
XX PN
XX PN (HOLM/) HOLMS R D.
XX PN
XX PN Holms RD;
XX PN
XX PN WPI: 2001-294297/41
XX PN
XX PN Novel regulatory or unfolding peptides of ezrin that binds to
XX PN heprecceptor, useful for inducing immune response for treating
XX PN infectious diseases and cancer
XX PS Claim 26: Page 37: 4app: English.
XX PS
XX CC the heprecceptor is a novel active site in human ezrin. Ezrin regulates
XX CC the structure of the cortical cytoskeleton to control cell surface
XX CC topography. The present invention relates to peptides (see AAB82021 to
XX CC AAB82041) that bind to heprecceptor with greater affinity than HEPI (see
XX CC AAB82046). The heprecceptor binding peptides are useful for inducing
XX CC immune response, and for treating infectious diseases, cancer and
XX CC HIV-related dementia. The present peptide binds to domain A of the
XX CC heprecceptor (AAB82019).
XX SQ Sequence 11 AA:

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Best Local Similarity 100.0%; Pred. No. 21;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MLRLQ 5
| | | | |

Db 6 MLRLQ 10

RESULT 4

AA882038
ID AA882038 standard; peptide; 12 AA.

XX AAB82038;

AC AAB82038;

XX 13-JUN-2001 (first entry)

XX Human hepreceptor domain A binding peptide kupe2132.

XX Human; hepreceptor, cytostatic, anti HIV, antiobiotic;

KW neotropic; immune response inducer; ezrin; infectious diseases; cancer;

KW HIV-related dementia.

XX Homo sapiens.

XX Key

XX Modified-site ID Location/Qualifiers

FT /note "Optionally phosphorylated"

XX GN2354241 A.

XX 21-MAR-2001.

XX 17-SEP-1999; 99CH-0021881.

XX 17-SEP-1999; 99CH-0021881.

XX (HOLMS) HOLMS R. D.

XX Holms RD;

XX WPI: 2001 293287/31

XX Novel regulatory or stabilizing peptides of ezrin that binds to

PT hepreceptor, useful for inducing immune response for treating

PT infectious diseases and cancer.

XX Claim 24: Page 36; 42pp; English.

XX The hepreceptor is a novel active site in human ezrin. Ezrin regulates

CC the structure of the cortical cytoskeleton to control cell surface

CC topology. The present invention relates to peptides. (see AAB82038 to

CC AAB82041) that bind to hepreceptor with greater affinity than HIV. (see

CC AAB82046). The hepreceptor binding peptides are useful for inducing

CC immune response, and for treating infectious diseases, cancer and

CC HIV-related dementia. The present peptide binds to domain A of the

CC hepreceptor (AAB82019).

XX Sequence 12 AA;

XX Query Match 100.0%; Score 23; DB 22; Length 12.

XX Best Local Similarity 100.0%; Pred. No. 21;

XX Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0.

QY 1 MLRLQ 5

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Db 4 MLRLQ 8

RESULT 5

AA081119
ID AA081119 standard; peptide; 12 AA.

XX AA081119;

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09-APR-2002 (first entry)

Integrin-antagonist peptide #26.

IgG Fc; anticoagulant; thrombolytic; cytostatic;

antiinflammatory; immunosuppressive; osteopathic; antagonist;

laminin; saw-scaled viper; echistatin; integrin; selectin; vinculin;

platelet aggregation; angiogenesis; tumour; inflammation;

autoimmune disease; rheumatoid arthritis; osteoporosis.

XX Synthetic.

XX W0200181377-A2.

XX 01-NOV-2001.

XX 23-APR-2001; 2001WO-0513069.

XX 21-APR-2000; 2000US-198919P.

XX 03-MAY-2000; 2000US-201394P.

XX (AMGEN) AMGEN INC.

XX Felipe U, Kohno T, Lacey DL, Boone TC;

XX WPI: 2002-052015708.

Composition comprising integrin or adhesion antagonistic peptide and vehicle, useful for treating or preventing platelet aggregation, has a longer half-life than free peptide.

Claim 11: Page 19; 68pp; English.

The invention relates to a composition comprising an integrin/adhesion antagonistic peptide (I) and a vehicle e.g. IgG Fc. The peptides are based on laminin or saw-scaled viper echistatin and target integrin, selectin or vinculin. Also included are compounds of formula (Ia) and their multimers (X¹)_a-P¹-(X²)_b where:

P¹ = FC domain;

X¹ and X² = -(L¹)₁-e-P¹, (L¹)₁-e-P¹-(L²)₁-P², (L¹)₁-e-P¹-(L²)₁-P²-(L³)₁-e-P³ or (L¹)₁-e-P¹-(L²)₁-P²-(L³)₁-e-P³-(L⁴)₁-P⁴;

P¹-P⁴ = same or different (I);

L¹-L⁴ = same or different linkers;

a = 0 or 1, provided at least one of a and b = 1,

a nucleic acid that encodes (Ia), an expression vector containing the pharmacologically active compound (B) by covalently linking at least one FC domain to at least one amino acid sequence of a selected randomized (C) and any of six admininistrated peptides (Ib). The compositions are used prophylactically and therapeutically in the same way as (I), e.g. to inhibit platelet aggregation or angiogenesis (tumours), or to treat inflammation and autoimmune diseases (e.g. rheumatoid arthritis) and many different forms of osteoporosis, also for diagnosis. Attaching the vehicle (especially FC domain) to (I) increases the half-life (three (I) are normally degraded very quickly in vivo). The present sequence is an antagonist peptide of the invention.

XX Sequence 12 AA;

XX Query Match 100.0%; Score 23; DB 23; Length 12;

XX Best Local Similarity 100.0%; Pred. No. 21;

XX Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MLRLQ 5

| | | | |

Db 6 MLRLQ 10

RESULT 6

ABB72970

ID ABB72970 standard; Peptide; 12 AA.

XX AC AAB72970;
 XX AC
 XX DT 05-APR-2002 (first entry)
 XX DE Integrin binding peptide SEQ ID NO:1081.
 XX
 XX Modified peptide; mimetic; Fe domain; fusion; immunoglobulin G; IgG;
 KW EPO; erythropoietin; TPO; tumour necrosis factor alpha inhibitor;
 KW TNF-alpha inhibitor; interleukin 1 antagonist; IL-1 antagonist; TMP;
 KW TPO mimetic peptide; EPO mimetic peptide; EMP; VEGF antagonist;
 KW MMP inhibitor; antiinflammatory; antitumour; immunosuppressive;
 KW cytostatic; antiinflammatory; antiarthritic; antidiabetic; ophthalmological;
 KW antianemic; anorectic; antifertility; haemostatic; dermatological;
 KW neuroprotective; inflammatory disease; autoimmune disease; tumour growth;
 KW cancer; rheumatoid arthritis; diabetic retinopathy; infertility; obesity;
 KW sleep disorder; neurological degenerative disease; anaemia;
 KW thrombocytopaenia; metastatic tumour; systemic lupus erythematosus;
 KW Fanconi's Syndrome.
 XX Homo sapiens.
 OS Synthetic.
 XX
 XX W0200183525-A2.
 XX
 XX 08 NOV-2001.
 XX
 XX 02 MAY-2001; 2001WO-0514710.
 XX
 XX 03 MAY-2000; 2000US-0563286.
 XX
 XX (AMGEN) AMGEN INC.
 XX
 XX Felipe U, Liu C, Cheetham JC, Boone TC, Gudas JM;
 XX WPI; 2002-130313/17.
 XX
 XX Novel vehicle-peptide molecule or its multimers useful for treating
 PT inflammatory and autoimmune diseases, cancer, rheumatoid arthritis,
 PT diabetic retinopathy, obesity, sleep disorders and infertility.
 XX
 XX Claim 39; Page 47; 176pp; English.
 XX
 XX The present invention describes a vehicle peptide molecule (i) or its
 CC multimers. (i) can have antiinflammatory, antitumour, immunosuppressive,
 CC cytostatic, antirheumatic, antiarthritic, antidiabetic, ophthalmological,
 CC antianemic, anorectic, antifertility, haemostatic, dermatological and
 CC neuroprotective activities. (i) can be used as a therapeutic or
 CC prophylactic agent as well as for screening purposes. (i) is useful for
 CC diagnosing diseases characterised by dysfunction of their associated
 CC protein of interest, for identifying normal or abnormal proteins of
 CC interest, as a part of diagnostic kit to detect the presence of their
 CC proteins of interest in a biological sample. Additionally, (i) is useful
 CC for treating inflammatory and autoimmune diseases, tumour growth, cancer,
 CC rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders,
 CC infertility, and neurological degenerative diseases (i), comprising
 CC EPO mimetic compounds are useful for treating disorders characterised by
 CC low red blood cell levels such as anaemia. The EPO-mimetic comprising
 CC compounds are useful for treating conditions that involve an existing
 CC megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet
 CC deficiency, such as thrombocytopaenia, aplastic anaemia, metastatic
 CC tumour which result in thrombocytopaenia, systemic lupus erythematosus,
 CC and Fanconi's syndrome. ABB72403 to ABB74426 and ABL35695 to ABL35777
 CC represent amino acid and nucleic acid sequences used in the
 CC exemplification of the present invention.
 XX
 XX Sequence 12 AA;
 SQ

Query Match 100.0%; Score 23; DB 23; Length 12;
 Best Local Similarity 100.0%; Pred. No. 21;
 Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MURIO 5

DB 6 MURIO 10
 RESULT 7
 AAB82037
 ID AAB82037 standard; peptide; 13 AA.
 XX
 XX AAB82037;
 AC
 XX DT 13-JUN-2001 (first entry)
 XX
 XX Human hepreceptor domain A binding peptide Rupe2042.
 DE
 KW Human; hepreceptor; cytostatic; anti-HIV; antibiotic;
 KW acotropic; immune response inducer; ezrin; infectious diseases; cancer;
 KW HIV-related dementia.
 XX
 XX Homo sapiens.
 OS
 XX Key Location/Qualifiers
 FH Modified-site 11
 FT /note- "optionally phosphorylated"
 FT
 XX GH2354241-A.
 PN
 XX 21-MAR-2001.
 PD
 XX 17-SEP-1999; 99GB-0021881.
 PF
 XX 17-SEP-1999; 99GB-0021881.
 PR
 XX (HOLM/) HOLMS R D.
 PA
 XX Holms RD;
 PI
 XX WPI; 2001-293287/31.
 DR
 XX Novel regulatory or unfolding peptides of ezrin that binds to
 PT hepreceptor, useful for inducing immune response for treating
 PT infectious diseases and cancer
 XX
 XX Claim 22; Page 36; 42pp; English.
 PS
 XX The hepreceptor is a novel active site in human ezrin. Ezrin regulates
 CC the structure of the cortical cytoskeleton to control cell surface
 CC topography. The present invention relates to peptides (see AAB82021 to
 CC AAB82041) that bind to hepreceptor with greater affinity than HEPI (see
 CC AAB82046). The hepreceptor binding peptides are useful for inducing
 CC immune response, and for treating infectious diseases, cancer and
 CC HIV-related dementia. The present peptide binds to domain A of the
 CC hepreceptor (AAB82019).
 XX
 XX Sequence 13 AA;
 SQ
 Query Match 100.0%; Score 23; DB 22; Length 13;
 Best Local Similarity 100.0%; Pred. No. 23;
 Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 MURIO 5
 DB 5 MURIO 9
 RESULT 8
 AAB38185
 ID AAB38185 standard; Protein; 15 AA.
 XX
 XX AAB38185,
 AC
 XX DT 30 JAN 2001 (first entry)
 XX
 XX Human secreted protein sequence encoded by gene 9 SEQ ID NO:124.
 DE

DR WPI: 2001 183426/52.
 XX N-PSDB: AAK59784.
 PT Nucleic acids encoding human immune/hematopoietic antigen polypeptides,
 PT useful for preventing, diagnosing and/or treating cancers and
 PT metastasis.
 PS Claim 11: SEQ ID NO 14596, 3071pp + Sequence Listing, English.
 XX AAK54951 to AAK54702 encode the human immune/hematopoietic antigen (I)
 CC amino acid sequences given in AAK52170 to AAK51922. (I) have cytostatic
 CC activity, and can be used in gene therapy and vaccine production. (I)
 CC proteins and polynucleotides may be used in the prevention, diagnosis and
 CC treatment of diseases associated with inappropriate (I) expression. For
 CC example, they may be used to treat disorders associated with decreased
 CC expression by rectifying mutations or deletions in a patient's genome
 CC that affect the activity of (I) by expressing inactive proteins or to
 CC supplement the patient's own production of (I). Additionally, (I)
 CC polynucleotides may be used to produce the secreted (I), by inserting
 CC the nucleic acids into a host cell and culturing the cell to express the
 CC protein. (I) proteins and polynucleotides may be used to prevent,
 CC diagnose and treat immune/hematopoietic-related diseases, especially
 CC cancers and cancer metastases of hematopoietic-derived cells. AAK54703
 CC to AAK87694 represent human immune/hematopoietic antigen genomic
 CC sequences from the present invention. AAK54942 to AAK54949 and AAK52169
 CC represent sequences used in the exemplification of the present invention.
 XX
 SQ Sequence 57 AA;

Query Match 100.0%; Score 23; DB 22; Length 57;
 Best Local Similarity 100.0%; Pred. No. 95;
 Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MLRLQ 5
 II III
 DB 7 MLRLQ 11

RESULT 11

ID AA006711 standard; Protein: 73 AA.

AC AAC06711;

DT 06-NOV-2001 (first entry)

XX Human polypeptide SEQ ID NO 20603.

DE Human, cytokine, cell proliferation, cell differentiation; gene therapy;
 KW vaccine, peptide therapy, stem cell growth factor, hematopoiesis;
 KW Tissue growth factor; immunomodulatory; cancer; leukaemia;
 KW nervous system disorders; arthritis; inflammation.

XX Homo sapiens.

XX W02001164835-A2

XX 07-SEP-2001.

XX 26-FEB-2001; 2001WO-0504927

XX 28-FEB-2000; 2000US-051526

XX 18-MAY-2000; 2000US-0577400

XX (HYSE-) HYSEQ INC.

XX Tang YT, Liu C, Drmanac RT;

XX WPI: 2001-514848/56

XX N-PSDB: AAI86642.

XX Isolated nucleic acids and polypeptides, useful for preventing
 PT diagnosing and treating e.g. leukaemia, inflammation and immune

PT disorders -
 XX
 PS Claim 20: SEQ ID NO 20603; 1599pp + Sequence Listing; English.
 XX The invention relates to human polynucleotides (AA170041-AA193841) and
 CC the encoded proteins (AA00010-AA013910) that exhibit activity relating to
 CC cytokine, cell proliferation or cell differentiation or which may induce
 CC production of other cytokines in other cell populations. The
 CC polynucleotides and polypeptides are useful in gene therapy, vaccines or
 CC peptide therapy. The polypeptides have various cytokine-like activities,
 CC e.g. stem cell growth, factor activity, hematopoiesis regulating
 CC activity, tissue growth factor activity, immunomodulatory activity and
 CC activity, inhibin activity and may be useful in the diagnosis and/or
 CC treatment of cancer, leukaemia, nervous system disorders, arthritis and
 CC inflammation.
 CC Note: The sequence data for this patent did not form part of the printed
 CC specification, but was obtained in electronic format directly from WIPO
 CC at ftp.wipo.int/pub/published_pat_sequences.

XX SQ Sequence 73 AA;

Query Match 100.0%; Score 24; DB 22; Length 73;
 Best Local Similarity 100.0%; Pred. No. 125; 02;
 Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MLRLQ 5
 II III
 DB 68 MLRLQ 72

RESULT 12

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AC ABB10135;

DT 10-JAN-2002 (first entry)

XX Human cDNA SEQ ID NO: 443.

DE Human; gene therapy; neural disorder; immune system disorder;
 KW muscular disorder; reproductive disorder; gastrointestinal disorder;
 KW pulmonary disorder; cardiovascular disorder; renal disorder;
 KW proliferative disorder; inflammation.

XX Homo sapiens.

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XX 02-AUG-2001.

XX 17-JAN-2001; 2001WO-0501349.

XX 31-JAN-2000; 2000US-179055P.

XX 04-FEB-2000; 2000US-180628P.

XX 24-FEB-2000; 2000US-181664P.

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XX 16-MAR-2000; 2000US-189874P.

XX 17-MAR-2000; 2000US-190076P.

XX 18-APR-2000; 2000US-198124P.

XX 19-MAY-2000; 2000US-205515P.

XX 07-JUN-2000; 2000US-209457P.

XX 28-JUN-2000; 2000US-214886P.

XX 30-JUN-2000; 2000US-215135P.

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XX 07-JUL-2000; 2000US-216880P.

XX 11-JUL-2000; 2000US-217487P.

XX 11-JUL-2000; 2000US-217496P.

XX 14-JUL-2000; 2000US-218290P.

XX 26-JUL-2000; 2000US-220963P.

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